Amendments to the Claims

This listing of claims will replace all prior versions, and listings, of claims in the application:

- 1. (CURRENTLY AMENDED) A Gram-negative bacterium An Agrobacterium comprising an inducible regulatory sequence operatively linked to a nucleotide sequence encoding a encoding levansucrase which are contained within the genome of said Gram-negative bacterium Agrobacterium.
- 2. (CURRENTLY AMENDED) A Gram-negative bacterium An Agrobacterium comprising a recombinant nucleotide sequence containing an inducible regulatory sequence other than sacR operatively linked to a nucleotide sequence encoding a encoding levansucrase.
- 3. (CURRENTLY AMENDED) The Gram-negative bacterium The Agrobacterium of claim 1, wherein said nucleotide sequence encoding a encoding levansucrase is a sacB open reading frame.
 - 4. (CANCELED)
- 5. (CURRENTLY AMENDED) The Gram-negative bacterium The Argobacterium of claim 3, wherein said bacterium is Agrobacterium tumefaciens.
- 6. (CURRENTLY AMENDED) The Gram-negative bacterium The Argobacterium of claim 1, wherein said regulatory sequence comprises the *E. coli* lactose operon (SEQ. ID. NO. 1).
- 7. (CURRENTLY AMENDED) The Gram-negative bacterium The

 <u>Agrobacterium</u> of claim 1, wherein said regulatory sequence comprises the Pi 2(noc)

 promoter (SEQ. ID. NO. 2) and noc 1 operon (SEQ. ID. NO. 2).
- 8. (CURRENTLY AMENDED) The Gram negative bacterium The

 Agrobacterium of claim 1, wherein said regulatory sequence comprises the P_{BAD}

 promoter (SEQ. ID. NO. 5) and araC cis element (SEQ. ID. NO. 4).
 - 9.-33. (CANCELED)
- 34. (CURRENTLY AMENDED) The Gram-negative bacterium The

 Agrobacterium of claim 1, wherein the regulatory sequence comprises the traCDG

promoter (SEQ. ID. NO. 7) and the occ promoter (SEQ. ID. NO. 8).

35. (CURRENTLY AMENDED) The Gram-negative bacterium The Agrobacterium of claim 3, wherein the nucleotide sequence encoding a encoding levansucrase contains a second copy of a sacB open reading frame.

36.-43 (CANCELED)